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Pl^o of BAG-1 protein in a sample of body fluid containing breast cancer cells.--

REMARKS

Claims 11-14, 16, 20-27, 32-34, 36, 37, 44 and 50-61 are pending in the present application. Claims 11, 13, 14, 20, 21, 26, 32, 36, 37, 44, 50-61 have been amended herein. Claims 62-66 have been added herein. Following entry of the amendments, claims 11-14, 16, 20-27, 32-34, 36, 37, 44 and 50-61 will remain pending.

Regarding the Amendments

The claim amendments are supported by the specification and introduce no new matter. Specifically, claims 11-14, 16, 20-27, 32, 34, 36, 37, and 50-61 have been amended to clarify antecedent basis, as described herein below. Claim 44 has been amended to more explicitly link the purpose of the method, as recited in the preamble, to results obtained by performing the method. Specifically, claim 44 now recites in the claim body that the method results in "determining a lower risk of tumor recurrence or spread in the first group of patients suffering from breast cancer." Attached as Appendix A is a marked up version of the claim amendments in which text to be deleted is enclosed in brackets and text to be added is underlined.

New claims 62-66 are directed to methods for prognosis of disease-free or overall survival of an individual having a

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breast cancer tumor; for predicting the risk of tumor recurrence or spread in an individual having a breast cancer tumor; for determining the proper course of treatment for a patient suffering from breast cancer; for screening a breast cancer patient to determine the risk of tumor metastasis or chance of survival; or for determining risk of tumor recurrence or spread in a patient suffering from breast cancer, respectively, in which the level of BAG-1 protein expression is determined by measuring the level of BAG-1 protein in a sample of body fluid containing breast cancer cells.

As set forth above, the claim amendments do not add new matter. Therefore, Applicant respectfully requests that the Examiner enter the amendments.

Rejection Under 35 U.S.C. § 112, First Paragraph

The objection to the specification and corresponding rejection of claims 11-14, 16, 20-27, 32-34, 36, 37, 44, and 50-61 under 35 U.S.C. 112, first paragraph, as allegedly lacking enablement is respectfully traversed. Applicant submits that the full scope of these claims is enabled by the specification.

Regarding claims 11-14, 16, 20-24, 27, 32-34, 36, 37, 50, 52-57, 59 and 60, the Office Action alleges that the specification lacks enablement for methods in which BAG-1 protein expression levels are measured in a body fluid. Applicant respectfully submits that those skilled in the art would have been able to practice the claimed methods using a sample that is

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body fluid in much the same way as when using a tumor sample. In this regard, the specification provides teachings that a body fluid is a suitable sample for use in the claimed methods; a rationale for why a body fluid is a suitable sample; provides procedures for performing steps of the claimed methods; and working examples of measuring a level of BAG-1 protein expression and correlating levels of BAG-1 protein expression with clinical outcome. In view of these teachings in the specification, Applicant submits that one skilled in the art would have been able to use the claimed methods without undue experimentation.

First, the specification teaches that a body fluid is a suitable sample for use in the prognostic methods of the invention. For example, the specification teaches that "a body fluid sample, such as a lymph, blood or serum sample, or an exudate fluid sample such as the cancerous organ exudate (e.g., exudate from the breast) may be collected and used as the sample to be analyzed (page 8, lines 9-24)." In view of this teaching, those skilled in the art would have understood that the claimed methods can be practiced using a body fluid.

Second, the specification provides rationales as to why a body fluid would contain BAG-1, and why a level of BAG-1 protein expression in a body fluid would correlate with clinical outcome. In particular, the specification states that, "while BAG levels will typically be measured within the cancerous cells of a patient, levels of BAG will also be measured in a body fluid sample (e.g., serum) as a result of BAG having been secreted or otherwise released from cells (e.g., by cell rupture) (page 8,

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lines 18-23)," and indicates that cancerous cells of a patient "may be drawn from a body fluid such as lymphatic fluid, blood, serum, or a distally infected organ or exudate thereof" in the case of metastatic cancer (page 8, lines 15-18). Knowing why a body fluid would contain BAG-1 and that an increased level of BAG-1 can be observed in breast cancer patients, those skilled in the art would have understood that BAG-1 protein expression levels are useful prognostic determinants in the claimed methods.

Moreover, it was well known at the time of filing the present application that a body fluid, such as an exudate fluid sample, can contain cells that originate from a cancerous tumor. In particular, as described in Sauter et al. British Journal of Cancer 76(4):494-501 (1997), submitted herewith as Exhibit 1, it was well known that breast exudate fluid (nipple aspirate fluid or NAF) contains exfoliated breast epithelial cells, which can include breast cancer cells because breast cancer develops from ductal and lobular epithelium (page 498, first paragraph in "Discussion" section). As such, it would be expected that cells originating from a breast cancer tumor and contained in a body fluid would be used in the same manner as a sample from the breast cancer tumor in the claimed methods.

Third, it was routine at the time of filing the present application to obtain body fluid samples from patients, and to use antibody-based methods to detect a level of protein expression. Evidence that body fluid sample collection was routine at the time of filing this application, is found for example in Sauter et al. This article describes routine methods

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for collecting breast exudate fluid (page 495, "Aspiration technique") and for determining if a biomarker correlates with breast cancer (page 497, "Statistical Analysis"). In view of the routine nature of body fluid sample collection and statistical analysis for correlating a biomarker with breast cancer risk at the time of filing the present application, Applicant submits that undue experimentation would not have been required to practice the claimed methods using a body fluid.

Moreover, the specification provides guidance for using BAG-1 specific antibody to determine BAG-1 protein expression levels. In particular, the specification describes that detection of BAG protein expression levels can be carried out using well-known methods such as histochemical staining, Western Blot analysis, immunoprecipitation and immunoassays, such as an ELISA, immuno-PCR (page 13, lines 21-29). This guidance would have been used by those skilled in the art for practicing the claimed methods using a body fluid sample.

Fourth, the specification provides description and working examples of correlating survival of an individual having a breast cancer tumor with levels of BAG-1 protein expression in a tumor sample, which would have been used by one skilled in the art when practicing a method of the invention using a body fluid. For example, the specification provides working examples of procedures for determining a level of BAG-1 protein expression that correlates with disease-free or overall survival (page 35, line 5, to page 36, line 6); procedures for classifying patients having high levels of expression of BAG-1 protein relative to a

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reference level; and procedures for identifying groups of patients having low or high levels of BAG-1 protein expression (page 33, line 25, to page 34, line 16). Applicant submits that those skilled in the art would have been guided by these teachings in the specification when practicing a method of the invention using a body fluid sample.

Finally, considering the high level of skill in the art surrounding diagnostics development at the time of filing the present application and in view of the above-described ample amount of direction provided in the specification, Applicant submits that undue experimentation would not have been required to practice the full scope of the claimed invention. Accordingly, Applicant submits that the full scope of claims 11-14, 16, 20-27, 32-34, 36, 37, 44, and 50-61 is enabled by the specification, and requests removal of the rejection of these claims under 35 U.S.C. 112, first paragraph.

Regarding claims 25, 26, 51, 58, 44 and 61

The Office Action states that the rejection of claims 25, 26, 44, 51, 58 and 61 is maintained, in part, because the specification allegedly fails to enable methods where measurement of BAG-1 protein expression is correlated with risk of tumor recurrence or spread. The Office Action asserts that tumor recurrence or spread do not necessarily predict disease-free or overall survival, but rather are merely contributing factors to the measurement of disease-free or overall survival. Applicant respectfully disagrees with the notion that tumor recurrence or

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spread are not directly correlated with disease-free or overall survival.

In contrast with these statements in the Office Action, the specification teaches that "disease-free survival," means a lack of tumor recurrence and/or spread, as well as the fate of a patient after diagnosis, with respect to the effects of the cancer on the life-span of the patient. This teaching remains valid today as is noted, for example, in an American Cancer Society website reference article entitled "What is Metastatic Cancer?" (submitted herewith as Exhibit 2). The article states that most people who die of cancer have metastasis (also referred to in the article as recurrence) at the time of their death, and that such metastasis "are directly responsible for the majority of cancer deaths" (page 2, first paragraph under heading "Significance of Metastasis to the Patient").

Thus, it would have been understood by those skilled in the art that risk of tumor recurrence or spread in a cancer patient correlates with disease-free or overall survival of that patient. Because the specification teaches that disease-free survival means a lack of tumor recurrence and/or spread; and demonstrates that levels of BAG-1 protein expression correlate with disease-free or overall survival of an individual having a breast cancer tumor, those skilled in the art would have understood that levels of BAG-1 protein expression are correlated with risk of tumor recurrence or spread. In view of the above, Applicant submits that claims 25, 26, 51, 58, 44 and 61 are enabled by the specification.

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Regarding claims 34, 36, 37, 53 and 60

The Office Action states that the rejection of claims 34, 36, 37, 53 and 60 is maintained, in part, because the specification allegedly fails to teach which levels of BAG-1 protein expression allow the division into two groups and how to determine the cut-off values of BAG-1 that would be necessary to divide a group of patients into two groups.

Applicant submits that the specification exemplifies how to determine cut-off values of BAG-1 protein expression useful for dividing a group of patients into two groups, and in doing so, teaches how to determine which levels of BAG-1 protein expression can be used to divide a group of patients into two groups. In particular, the specification teaches, on page 34, lines 3-13, that:

To set cut-offs for dichotomization of data into groups having high and low levels of expression, the H-score data for the entire data-set were displayed as dot-histograms with H-score on the x-axis and the number of patient samples having a given H-score on the y-axis. An H-score ≥ 150 was determined by this approach to be appropriate for use as a cut-off for high levels of BAG-1 expression. H-score data were displayed as histograms and dichotomized into high versus low expression level groups using optimized cut-offs

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for BAG-1 (H-score \geq 150), Bcl-2 (H-score \geq 180), and
Bax (H-score \geq 140).

Moreover, Applicant submits that the level of skill in the art surrounding diagnostics development was advanced at the time of filing the present application. For example, data analysis and dichotomization using statistical tools such as Kaplan-Meier analysis and Cox-regression analysis was routine. In addition, methods for immunoscoreing, such as those described in King et al., 1985, Cancer Research 45: 293-296, as referenced in the specification on page 34, lines 15-16, were used routinely.

In view of the above teachings in the specification, and what was known in the art, Applicant submits that those skilled in the art would have been able to determine cut-off values of BAG-1 protein expression useful for dividing a group of patients into two groups without undue experimentation. Therefore, Applicant requests removal of the rejection of claims 34, 36, 37, 53 and 60.

Rejection under 35 U.S.C. § 112, second paragraph

The rejection of claims 11-14, 20, 21, 26, 32, 33, 36, 37, 44, 50-53, and 55-61 under 35 U.S.C. § 112, second paragraph, as allegedly indefinite is respectfully traversed.

The Office Action states that the rejected claims are indefinite because they lack antecedent basis for the phrases

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"the level of BAG-1 expression," "the amount of BAG-1 protein," and "said level of BAG-1 expression."

Applicant respectfully submits that the scope of the rejected claims would be reasonable ascertainable by those skilled in the art because "the level of BAG-1 expression" as recited in the dependent claims would have been understood to refer to "the level of BAG-1 protein expression" as recited in the independent claims. Similarly, the "amount of BAG-1 protein" would have been understood to refer to "level of expression of BAG-1 protein." Nevertheless, Applicant has amended several claims to clarify antecedent basis, as described below.

Claims 11, 13, 14, 20, 21, 50, 55, 56 and 57 have been amended to recite a "level of BAG-1 protein expression" to clarify antecedent basis with respect to claim 16. Therefore, Applicant respectfully requests removal of this rejection of claims 11-14, 20, 21, 50, and 56.

Claims 26, 51 and 58 have been amended to recite a "level of BAG-1 protein expression" to clarify antecedent basis with respect to claim 25. Therefore, Applicant respectfully requests removal of this rejection of claims 26, 51 and 58.

Claim 32 has been amended to replace the term "the amount of BAG-1 protein" with "the level of expression of BAG-1 protein" to clarify antecedent basis with respect to claim 27. Therefore, Applicant respectfully requests removal of this rejection of claims 32 and 33.

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Claims 52 and 59 have been amended to replace the term "level of BAG-1 expression" with a "level of expression of BAG-1 protein" to clarify antecedent basis with respect to claim 27. Therefore, Applicant respectfully requests removal of this rejection of claims 52 and 59.

Claims 36, 37, 53 and 60 have been amended to recite a "level of BAG-1 protein expression" to clarify antecedent basis with respect to claim 34. Therefore, Applicant respectfully requests removal of this rejection of claims 36, 37, 53 and 60.

Claim 61 has been amended to recite a "level of expression of BAG-1 protein" to clarify antecedent basis with respect to claim 44.

The Office Action alleges that claim 44 is indefinite because it lacks a recitation of the correlation between performance of the method steps and the purpose of the method that is stated in the preamble.

Applicant respectfully submits that the scope of claim 44 would be reasonable ascertainable by those skilled in the art. In this regard, it would have been understood that if a first group has a lower likelihood of tumor recurrence or spread than the second group, it follows that the first group has a lower risk of tumor recurrence or spread. Nevertheless, Applicant has amended claim 44 to include the phrase "thereby determining risk of tumor recurrence or spread in a patient suffering from breast cancer" to recite a correlation between the

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preamble statement and performance of the method steps.
Therefore, Applicant respectfully requests removal of this
rejection of claim 44.

CONCLUSION

In light of the amendments and remarks herein,
Applicant submits that the claims are now in condition for
allowance and respectfully requests a notice to this effect.
Should the Examiner have any questions, she is invited to call
the undersigned agent or Cathryn Campbell.

Respectfully submitted,

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APPENDIX A

11. (Twice Amended) The method of claim 16, wherein said level of BAG-1 protein expression is determined by measuring the amount of BAG-1 protein product using an immunoassay.

13. (Amended) The method of claim 16, wherein said level of BAG-1 protein expression is determined prior to lymph node involvement of said cancer.

14. (Amended) The method of claim 16, wherein said level of BAG-1 protein expression is determined after lymph node involvement of said cancer.

20. (Amended) The method of claim 16, wherein said level of BAG-1 protein expression is determined by measuring the level of BAG-1 protein that is detectable in samples selected from the group consisting of breast tumor tissue, blood, serum, and plasma.

21. (Amended) The method of claim 16, further comprising determining if said level of BAG-1 protein expression represents an overproduction that is above a reference level of BAG-1 expression.

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26. (Amended) The method of claim 25, further comprising:

(a) determining an overproduction level for BAG-1 protein, said level being in excess of a minimum amount statistically determined to be indicative of decreased likelihood of tumor recurrence or spread;

(b) determining the level of BAG-1 protein expression in said tumor sample; and

(c) predicting said risk of tumor recurrence or spread wherein an overproduction level of BAG-1 protein in the tumor sample is negatively associated with the likelihood of tumor recurrence or spread.

32. (Twice Amended) The method of claim 27, wherein the [amount of] level of expression of BAG-1 protein is measured using an immunoassay.

36. (Amended) The method of claim 34, wherein said level of BAG-1 protein expression is determined prior to lymph node involvement.

37. (Amended) The method of claim 34, wherein said level of BAG-1 protein expression is determined after lymph node involvement of said cancer.

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44. (Twice Amended) A method for determining risk of tumor recurrence or spread in a patient suffering from breast cancer, said method comprising:

(a) determining, using a BAG-1 specific antibody, the level of expression of BAG-1 protein in a cancerous tissue of a patient during stage I or stage II of said cancer; and

(b) classifying said patient as belonging either to a first group of patients having high levels of expression of BAG-1; or a second group of patients having low levels of expression of BAG-1,

wherein said first group has a lower likelihood of tumor recurrence or spread than said second group, thereby determining a lower risk of tumor recurrence or spread in the first group of patients suffering from breast cancer.

50. (Amended) The method of claim 16, wherein said level of BAG-1 protein expression is determined by measuring the level of BAG-1 protein in a sample of breast tumor tissue.

51. (Amended) The method of claim 25, wherein said level of BAG-1 protein expression is determined by measuring the level of BAG-1 protein in a sample of breast tumor tissue.

52. (Amended) The method of claim 27, wherein said level of expression of BAG-1 protein [expression] is determined by measuring the level of BAG-1 protein in a sample of breast tumor tissue.

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53. (Amended) The method of claim 34, wherein said level of BAG-1 protein expression is determined by measuring the level of BAG-1 protein in a sample of breast tumor tissue.

55. (Amended) The method of claim [49] 54, wherein said level of BAG-1 protein expression is determined by measuring the level of BAG-1 protein in a sample of breast tumor tissue.

56. (Amended) The method of claim 16, wherein said level of BAG-1 protein expression is determined by immunohistochemistry.

57. (Amended) The method of claim [50] 55, wherein said level of BAG-1 protein expression is determined by immunohistochemistry.

58. (Amended) The method of claim 25, wherein said level of expression of BAG-1 protein [expression] is determined by immunohistochemistry.

59. (Amended) The method of claim 27, wherein said level of BAG-1 protein expression is determined by immunohistochemistry.

60. (Amended) The method of claim 34, wherein said level of BAG-1 protein expression is determined by immunohistochemistry.

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61. (Amended) The method of claim 44, wherein said level of expression of BAG-1 protein [BAG-1 expression] is determined by immunohistochemistry.

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Nipple aspirate fluid: a promising non-invasive method to identify cellular markers of breast cancer risk

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Summary To evaluate the feasibility of nipple aspiration and to identify intermediate markers of breast cancer risk, nipple aspirate fluid (NAF) was collected from 177 subjects using a modified breast pump. The first 33 subjects demonstrated that we could obtain NAF quickly, reliably and repeatedly. Specimens from the remaining 144 subjects were collected to evaluate promising cellular biomarkers. NAF was obtained in 167 out of 177 (94%) subjects overall and in 99% of the 144 most recent subjects. Sufficient NAF was obtained to evaluate cytology in 160 out of 167 (96%) cases and specimens were sufficiently cellular to analyse DNA markers in 53% of cases. Among the last 144 subjects, menopausal status did not influence the ability to obtain NAF. NAF cytology correlated with increased breast cancer risk ($P = 0.002$). Using computerized image analysis of NAF epithelial cells, DNA index ($P = 0.0002$), percentage of cells in G₂M ($P = 0.05$) and percentage of cells with hypertetraploidy ($P = 0.002$) increased as cytology became more abnormal. Our data indicate that NAF can be obtained in essentially all eligible subjects; that breast epithelial cells are evaluable in > 95% of NAF samples for cytology and in over half of NAF samples for DNA index (ploidy) and cell cycle analysis; and that abnormal NAF cytology correlates with increased breast cancer risk. This suggests that biomarkers identified in nipple aspirate fluid may prove useful either as an adjunct to currently accepted breast cancer screening methods, or to evaluate response to a chemopreventive agent.

Keywords: nipple aspirate fluid; intermediate biomarker; breast cancer risk

Despite the death of 45 000 women annually in the United States from the disease, the ability to assess a woman's risk of developing breast cancer remains inadequate. The only well-established procedures to screen subjects for breast cancer are physical examination and mammography. Unfortunately, physical examination does not identify a significant number of early breast cancers and mammograms miss 10–40% of early breast cancers (Giuliano, 1994). Additional screening tools to identify precancer and early breast cancer are urgently needed.

Although the early detection of breast cancer will lead to a higher cure rate, the ideal form of treatment is prevention. The prevention of breast cancer is hindered by the difficulty in identifying an effective agent. Effective agents are difficult to identify in part because of the long period required for breast cancer to develop and, consequently, the requirement for lengthy clinical trials to test the efficacy of the agent, if the end point is the prevention of cancer. One way to shorten the time to finding of an effective agent is the identification of intermediate biomarkers, which are biological alterations in cells or tissue that occur between the time of initiation and tumour invasion. The theory is that an agent that partially or completely reverses the intermediate biomarker back to a normal phenotype may be interrupting carcinogenesis. Validation of the biomarker would require that the agent also decrease the incidence of cancer. Evaluating the effect of the agent requires the analysis of tissue, cells or non-cellular fluid. Nipple

aspiration, which is simple, quick, reliable and repeatable, would be an excellent tool to evaluate the efficacy of a chemopreventive agent should intermediate biomarkers be identified in the fluid.

Present efforts to evaluate the breast directly, through evaluation of either tissue or individual cells, have been hindered because the analysis of these specimens generally requires an invasive procedure. The adult non-pregnant, non-lactating breast secretes fluid into the breast ductal system. This fluid can be obtained through aspiration of the nipple with a modified breast pump. Refinements in the ability to obtain this fluid, as well as epidemiological studies to identify subjects most likely to yield NAF, have been on-going for over 20 years. Nipple aspiration has the attractiveness of quickly, painlessly, and non-invasively obtaining breast epithelial cells, the cells at risk for transformation to breast cancer. Nonetheless, limitations in the ability to obtain NAF in 30–50% of subjects has hindered the development of the technique as a tool for both breast cancer screening and evaluating response to treatment (Petrakis, 1993).

Our study had two goals. The first was to determine if the procedure was feasible. That is, could NAF be obtained consistently and were the NAF samples sufficiently cellular such that accepted markers of prognosis in breast tumours (e.g. cytology and ploidy) could be evaluated in the fluid? If it could be demonstrated that the procedure was feasible, the next goal was to identify intermediate biomarkers in the NAF that would correlate with breast cancer risk. Subjects of all risk categories were recruited. Cytological study of the Papanicolaou-stained smears and computerized image analysis were used to identify promising cellular markers of breast cancer risk. We evaluated markers such as cellular atypia, DNA index or ploidy and S-phase fraction, which in breast carcinoma tissue specimens are known to correlate with prognosis

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(Leis, 1991). We hypothesized that higher levels of these markers in NAF epithelial cells would correlate with increasing breast cancer risk. Because of a previous report indicating that NAF was more successful in premenopausal subjects, we also evaluated the influence of menopausal status on our ability to obtain NAF. Our findings form the basis of this report.

MATERIALS AND METHODS

Subjects

One hundred and seventy-seven non-Asian subjects aged 30–65 years (women undergoing mastectomy excepted) were recruited between August 1994 and June 1996 after approval of the Fox Chase Institutional Review Board. Asians, as well as women under 30 or greater than 65 years, were excluded because the success in obtaining NAF from the intact breast in these groups is known to be low (Petrukis, 1993). These subjects were categorized by their risk for breast cancer as having no risk factors, a first-degree relative with breast cancer, a history of curative treatment for ductal carcinoma in situ (DCIS) or invasive breast cancer, precancerous mastopathy (atypical hyperplasia (AH) or tubular carcinoma in situ (LCIS) or recently diagnosed DCIS) or recently diagnosed invasive cancer of the breast. We relaxed the age limit for women undergoing mastectomy to include subjects up to 72 years old because: (1) there are no data outlining the NAF yield from a breast that has just been removed and (2) our preliminary attempts to obtain NAF in mastectomy specimens from women up to 72 years of age were successful. Aspiration visits spaced either 3 days apart (subjects undergoing 6–12 aspirations) or 2 weeks apart (subjects undergoing two or three aspirations) were performed by a trained physician or nurse clinician.

Aspiration technique

After informed consent was obtained, nipple fluid was aspirated using a modified breast pump. The device consists of a 10-ml syringe attached to the end of a no. 4 endotracheal tube over which is placed a respiratory humidification adaptor. Each of these pieces is inexpensive and readily available in any hospital where mechanical respiratory support is provided.

For subjects in whom the intact breast was aspirated, the individual was seated in a comfortable position and the breast nipple cleansed with alcohol. After the alcohol evaporated, a warm, moist cloth was placed on each breast. After 1–2 min, the cloths were removed, the patient massaged her breast with both hands, and the plunger of the syringe was withdrawn to the 7-ml level and held for 15 s or until the patient experienced discomfort. A similar degree of suction was created with the syringe in mastectomy specimens obtained immediately after surgical excision. Fluid in the form of droplets was collected in capillary tubes. The quantity of fluid varied from 1 µl to 200 µl. The negative pressure produced was well tolerated. The procedure was repeated once on the same breast. Aspiration was then performed on the opposite breast, if present. If there was insufficient fluid from either breast to collect into one or more capillary tubes, or if the only breast that could be sampled (because of previous mastectomy) yielded no collected fluid, the patient was designated a non-secretor.

Occasionally, keratin plugs rather than NAF were obtained after suction was completed. The plugs were removed with an alcohol swab and suctioning repeated. Occasionally, suctioning was

performed two or three times to remove all of the plugs. Fluid was then obtained frequently. In order to obtain additional fluid, the nipple was gently compressed by the subject between her fingers. One or two additional droplets of fluid often appeared.

Cytology

Specimen preparation

The NAF was collected in 50-µl capillary tubes, rinsed into a container with 1 ml of 2% polyethylene glycol in ethanol-isopropanol and transported to the cytology laboratory for processing. In subjects with normal breast cancer risk, a family history of breast cancer, AH, or LCIS the samples were combined, given that the risk of invasive breast cancer is approximately equal in each breast. To evaluate the validity of this approach, early in our experience we collected NAF from nine subjects with AH or LCIS. The fluid from each breast was placed in separate containers. In seven out of nine cases, the cytology in both breasts was the same. In one of the remaining subjects, the abnormal NAF cytology was found in the breast opposite to where a prior biopsy had demonstrated AH, whereas in the second case the abnormal cytology was in the ipsilateral breast. If a cytology report was available from each breast, the report from the breast at highest risk was recorded. If a cytology report was available from each breast and the breasts were at equal risk, either the left or the right breast sample was selected randomly. If multiple cytological samples were analysed from the same breast, the most abnormal result was used in our analyses.

On the other hand, for subjects with previously treated DCIS or invasive breast cancer, only the breast without prior disease was aspirated. This was because the breast with disease had already been removed or radiated. For subjects with recently diagnosed but not yet definitively treated DCIS or invasive breast cancer, only the breast diagnosed with in situ or invasive disease was aspirated.

Table 1 Cytological criteria for nipple aspirate specimens

Scant mammary epithelial cells

This includes acellular specimens, those with only foam cells or with fewer than ten mammary epithelial cells

Benign mammary epithelial cells

This consists of specimens containing more than ten mammary epithelial cells without cytological atypia. This category encompasses normal mammary cells, apocrine metaplasia, and duct hyperplasia without atypia. We subdivide this category into:

- (a) Normal/non-papillary – single cells or small, loosely cohesive aggregates of cells.
- (b) Hyperplastic – large three-dimensional clusters of cells (indicative of duct hyperplasia or papillomatosis)

Atypical mammary epithelial cells

This category consists of specimens with ten or more mammary epithelial cells that exhibit atypical features. Atypia is defined as nuclear enlargement and/or irregularity, increased nuclear to cytoplasmic ratio or a chromatin distribution abnormally short of obvious malignant criteria. It is noted whether the atypical cells are single or in papillary clusters.

Malignant cells present

This category consists of specimens that contain cells with unequivocal criteria of malignancy. If the malignant cells are fewer than ten, the specimens are designated as 'scanty evidence'.

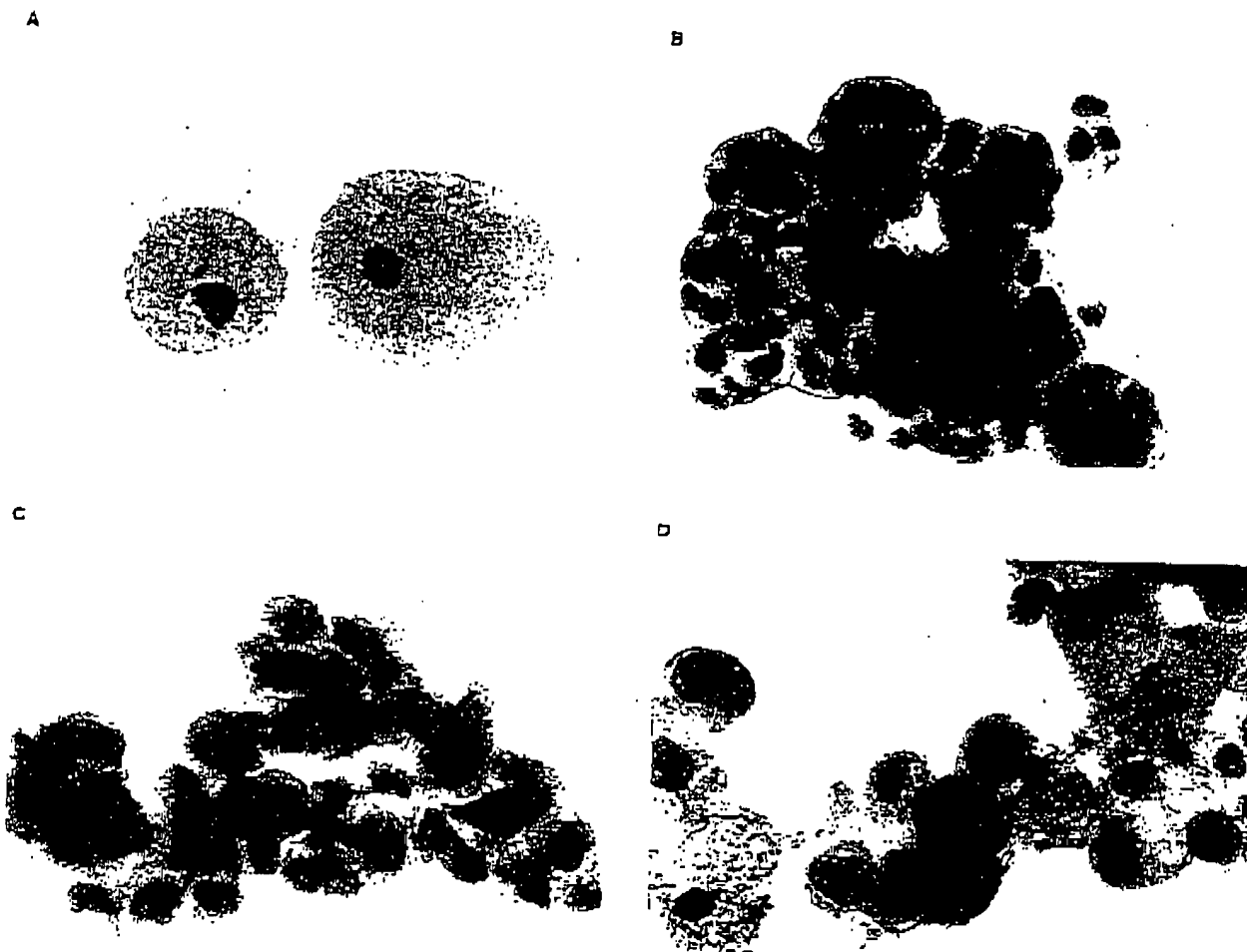


Figure 1 Photomicrographs of cytological preparations (Papanicolaou stain, $\times 1000$). (A) Foam cells, a frequent constituent of NAF; (B) papillary cluster of benign epithelial cells without atypia; (C) papillary cluster of atypical mammary epithelial cells; (D) malignant epithelial cells

In each case, the entire specimen was cytocentrifuged onto ten glass slides. Three of the slides were used for cytological examination. If the slides contained less than ten epithelial cells, two additional slides were examined. The remaining slides (five or seven) were stored for biomarker studies. The slides selected for cytological examination were washed twice in 95% ethanol for 5 min each, rehydrated in tap water and stained by the Papanicolaou method.

Specimen interpretation

The Papanicolaou-stained smears were examined by a cytopathologist experienced in breast cytology. Each specimen was designated (Table 1) as containing scant, benign, atypical or malignant cells, using a classification modified from King et al (1983). Representative photographs of various types of cells found in NAF, including benign, atypical and malignant epithelial cells, as well as foam cells, are illustrated in Figure 1.

Image analysis

Only specimens with adequate cellularity (cytology classes II–IV) were evaluated by image analysis. Although 88 specimens (53%)

were sufficiently cellular to perform image analysis, this technique was not employed until more than half of the subjects had been accrued. Thus far, 48 specimens have been evaluated.

Specimen preparation

A standardized quantitative DNA staining kit (RIAS Feulgen 5min Kit, Roche Image Analysis Systems, Elon College, NC, USA) was used following the manufacturer's instructions. In brief, after rehydration the slides were processed for hydrolysis with 5 N hydrochloric acid for 60 min and then transferred to the staining solution (Schiff's reagent) for 1 h, rinsed, dehydrated and mounted with synthetic resin.

Specimen interpretation

The Roche Pathology Workstation (Ellison et al, 1995) was used to evaluate nuclear ploidy, S-phase fraction, percentage of cells in G₂M and percentage hyperdiploid cells (cells with more than twice their complement of DNA), using computerized image analysis of cytocentrifuged nipple aspirate epithelial cells stained with a standard Feulgen preparation. Human lymphocytes were used as a control diploid cell population. All epithelial cells if

Table 2 Yield of fluid and epithelial cells by menopausal status

Status	Fluid obtained (%)		Cytology*			
	Subjects	Attempts*	Cellular	Low cellularity	No fluid	Non-cellular studies only
Overall						
Premenopausal	83/88 (97)	221/228 (97)	45	35	3	3
Post-menopausal	84/81 (92)	127/146 (87)	43	37	7	4
Totals	167/177 (94)	348/374 (93)	88	72	10	7
Last 144 subjects						
Premenopausal	70/71 (99)	198/200 (99)	38	29	1	3
Post-menopausal	73/73 (100)	107/111 (96)	40	29	0	4
Totals	143/144 (99)	305/311 (98)	78	58	1	7

*Cytology: if more than one cytological specimen was collected on a given subject, the most cellular specimen was counted; *Attempts: each subject may have undergone up to 12 aspiration attempts.

Table 3 Yield of fluid and epithelial cells by breast cancer risk category

Risk	Fluid obtained (%)		Cytology*			
	Subjects	Attempts*	Cellular	Low cellularity	No fluid	Non-cellular studies only
Normal risk or family history	48/49 (98)	217/220 (99)	19	29	1	0
Personal history of breast cancer	14/16 (88)	18/20 (90)	3	8	2	3
Precancerous mastopathy	32/39 (82)	40/51 (78)	22	10	7	0
Invasive cancer	73/73 (100)	73/73 (100)	44	25	0	4
Totals	167/177 (94)	348/374 (93)	88	72	10	7

*Cytology: a cellular specimen contains ≥ 10 breast epithelial cells on a slide. If more than one cytology specimen collected on a given subject, the more cellular specimen was counted; *attempts: each subject may have undergone one, two or three aspiration attempts; *personal history, precancerous mastopathy: the nine subjects in whom fluid was not obtained were enrolled very early in the study, before the standardization of our aspiration technique.

Table 4 Cytology in nipple aspirate fluid specimens by risk category

Risk	n	Cytological class				P-value*
		I	II	III	IV	
Normal risk or family history (FH)	50	29	15	6	0	NS
Personal history of breast cancer	10	8	2	0	0	
Precancerous mastopathy (PM)	31	10	11	8	2	
Invasive cancer (IC)	69	25	26	11	7	
Overall						0.002

*P-value: NS, not significant; 0.005, PM vs normal risk or family history of breast cancer; 0.004, IC vs normal risk or family history of breast cancer; 0.002, overall difference between the groups.

under 100 were present on a slide or a minimum of 100 cells if more were present were measured per case and type of stain. The average number of cells counted per specimen was 30.

Statistical analysis

Cumulative logistic regression models (Agresti, 1990) were fitted to the data to determine whether the distribution of cytological groupings was associated with the percentage of cells in G₂M, percentage of cells in S-phase, DNA index, current use of hormone replacement therapy, current use of oral contraceptives,

previous pregnancy, number of livebirths, number of miscarriages and whether the subject had difficulty becoming pregnant. Similarly, these methods were used to determine whether increasing breast cancer risk categories were associated with cytological grouping, percentage of hypertriploid cells, percentage of cells in G₂M, percentage of cells in S-phase and ploidy. The data did not support the proportional odds assumption for the relationship between cytological group and percentage hypertriploid cells. Therefore, the Kruskal-Wallis k-sample procedure (Mollander and Wolfe, 1973) was used to test the null hypothesis that individuals in the different cytological groups had identical distributions of hypertriploid cells in the nipple aspirate fluid. Failure to reject the null hypothesis ($P > 0.05$) indicates that subgroups could not be distinguished. Rejection of the null hypothesis ($P < 0.05$) indicates that there are at least two subgroups where the hypertriploid values are not identically distributed.

Fisher's exact two-sided procedures (Agresti, 1990) were used to determine whether a subject's ability to successfully provide NAF was associated with menopausal status. Logistic regression models were fitted to the data using generalized estimating equations methods (Liang and Zeger, 1986), to determine whether the menopausal status of the subject was associated with the probability of successfully obtaining NAF at any given attempt. To account for any potential within-subject correlation in the outcome variable, the test statistics were computed using the robust variance estimates.

Table 5 Cellular markers in nipple aspirate fluid specimens by cytological class

Cytology	Sample Size	DNA Index			G ₂ M			Hypertetraploid cells (%)		
		Mean	Median	Range	Mean	Median	Range	Mean	Median	Range
No atypia	28	1.06	1.03	0.74-1.59	1.51	0	0-5.67	2.14	0	0-26.79
Atypical hyperplasia	14	1.16	1.11	0.88-1.55	2.95	1.14	0-9.8	0.89	0	0-8.6
Malignant	6	1.77	1.78	1.31-2.35	9.20	3.85	0-37.8	17.29	15.79	0-45.16
P value				0.0002		0.05			0.002	

Because image analysis has only recently been employed, 48 specimens have been evaluated in both the DNA index and % hypertetraploid categories.

RESULTS

The correlation of breast cancer risk and NAF cytology with a variety of clinical factors, including current use of birth control pills (BCP), current use of hormone replacement therapy (HRT), whether the subject was ever pregnant, the number of livebirths, the number of miscarriages, the subject's difficulty becoming pregnant, the phase of the menstrual cycle in which the sample was obtained and the ease with which NAF was obtained were evaluated. None of these factors correlated with breast cancer risk or NAF cytology, although the power of the analysis of current use of birth control pills (BCP) and current use of hormone replacement therapy (HRT) was diminished by having an unbalanced population (few subjects were currently using either BCP or HRT). None of the clinical factors analysed affected the ability to obtain NAF. Nor did the interval between aspirations or the number of aspirations affect the ability to obtain fluid or the volume of NAF obtained on a given visit. Our cytologist did not identify blood in any NAF sample from women with an intact breast and rarely found blood in NAF from mastectomy specimens. Using image analysis, the amount of DNA in the breast epithelial cells obtained from NAF and the percentage of cells in each phase of the cell cycle were also evaluated. Although ploidy, the percentage of hypertetraploid cells and the percentage of cells in G₂M did correlate with NAF cytology, the percentage of cells in the S-phase of the cell cycle did not.

To evaluate the influence of menopausal status on our ability to obtain NAF, our success in premenopausal vs post-menopausal subjects was compared (Table 2). The data were presented in two ways: first as the entire experience with 177 subjects and then with our most recent 144 subjects, which reflects our success with nipple aspiration once the technique was standardized. We had greater overall success per aspiration in pre- vs post-menopausal subjects (97% vs 87%, $P < 0.001$), although this difference was not seen in the last 144 subjects (98% vs 96%, P not significant).

Early in the study, we also assessed the effect of breast irradiation on our ability to obtain NAF specimens. Four subjects with DCIS who underwent prior radiation had their radiated breast aspirated twice or three times for a total of 11 aspirations. NAF was obtained only twice and both samples had scant cellularity.

Our success in obtaining both NAF and cellular specimens in subjects classified by breast cancer risk is outlined in Table 3. NAF was obtained in 167 out of 177 subjects in 348 out of 374 attempts. Fifty-three per cent (88 out of 167) of the specimens contained ten or more breast epithelial cells on a slide, the criterion used to carry out cellular biomarker determinations.

The correlation of nipple aspirate cytology and image analysis parameters (DNA index, percentage of hypertetraploid cells, S-phase fraction, and percentage of cells in G₂M) with risk category

was next evaluated (Table 4). Risk was classified as normal risk or family history of breast cancer, personal history of breast cancer (prior DCIS or invasive cancer), precancerous mastopathy or invasive cancer. Overall, nipple aspirate cytology ($P = 0.002$) was significantly associated with breast cancer risk. Each increased risk category was compared with normal risk/family history of breast cancer. Pairwise comparisons revealed that samples from individuals with invasive cancer ($P = 0.004$) or precancerous mastopathy ($P = 0.005$) were significantly more likely to contain more abnormal cells.

Cellular markers, including DNA index, percentage of cells in the S-phase fraction, percentage of cells in G₂M of the cell cycle and the percentage cells with more than twice the normal amount of DNA (percentage hypertetraploid cells) were evaluated using image analysis (Tables). DNA index ($P = 0.0002$), the percentage of cells in G₂M ($P = 0.05$) and the percentage of hypertetraploid cells ($P = 0.002$) increased as cytology became more abnormal (Figure 2). Although there was a gradual increase in the DNA index and in the percentage of cells in G₂M of the cell cycle that gradually increased as the cytology became more abnormal, the pattern seen for cells with hypertetraploidy was slightly different. In the latter case, the percentage of hypertetraploid cells was similar for cytological classes II and III, whereas there was a sudden increase in the number of hypertetraploid cells with class IV (malignant) cytology.

DISCUSSION

The breast ducts of adult non-pregnant women secrete small amounts of fluid (Keynes, 1923). This fluid does not escape because the nipple ducts are occluded by smooth muscle contraction, dried secretions and keratinized epithelium. Breast fluid can be obtained by nipple aspiration in a significant proportion of women without spontaneous nipple discharge with the use of a modified breast pump (Petrakis et al, 1975). This fluid contains several types of cells, including exfoliated breast epithelial cells (King et al, 1975). Because breast cancer develops from ductal and lobular epithelium, NAF is a potentially useful epidemiological and clinical research tool. A major limitation of the technique has been the lack of ability to obtain NAF in all women, and when fluid was obtained it frequently contained few or no breast epithelial cells.

Published data from Petrakis et al (1993) indicate that the highest yield of NAF is from women aged 30-55 years with early onset of menarche, non-Asian ethnicity and prior parity and/or lactation. This procedure has yielded NAF in over 50% of premenopausal Caucasian and African-American women over the age of 30. With a definition of ≥ 10 epithelial cells as an adequate aspiration, others have obtained an adequate specimen in 42% of women aspirated (King et al, 1983). The nipple aspiration

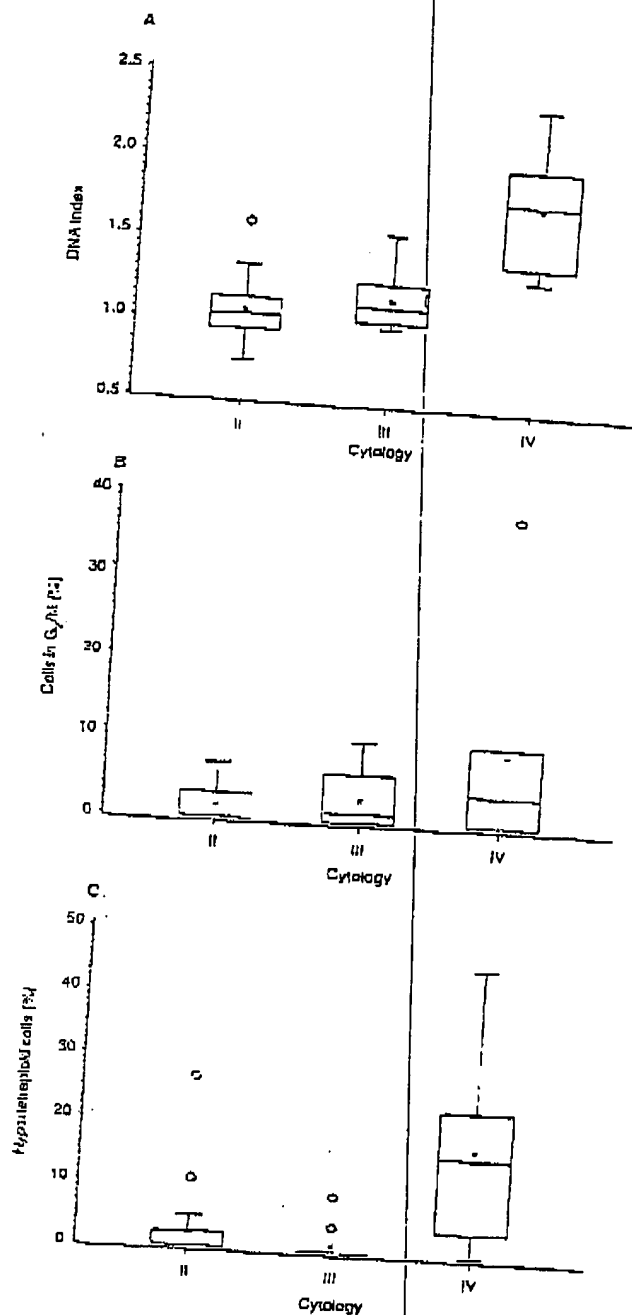


Figure 2. Box plots of image analysis markers, indicating the median (•), mean (horizontal line within the box), 25–75% confidence intervals (outer box horizontal lines), and standard deviation (bars above and below the boxes). (A) DNA Index as a function of cytological class; (B) percentage cells in G₂/M, as a function of cytological class; (C) percentage hyperdiploid cells, as a function of cytological class.

technique has been employed without significant side-effects in over 7000 women.

We were able to improve our success in obtaining NAF from 43% at the start of the study to 99% at the present time. Early in the study, we found that we could often obtain NAF at a second or

third visit, even though NAF was not obtained at the first visit. This ability to obtain NAF at some but not all visits was not related to the phase of the menstrual cycle in which the aspiration was performed. Now we are able to obtain NAF in over 95% of aspiration attempts. In the few subjects who did not yield NAF at the first visit, we were always able to obtain NAF during subsequent visits if they were willing to return.

Although Petrakis et al (1993) were more successful in obtaining NAF in pre- than in post-menopausal subjects, they also found that age (30–55 years) and ethnicity (non-Asian) were more important predictors of success in obtaining NAF than menopausal status. Our interest in the possible influence of menopausal status on NAF yield led us to determine its importance in our 177 subjects. Although early in our experience we had greater success (97% vs 87%) in obtaining NAF from premenopausal subjects (Table 2), menopausal status did not influence our ability to obtain NAF (99% for pre- vs 96% for post-menopausal subjects) from our last 144 subjects.

In this report, we demonstrate a higher success rate in obtaining NAF than has generally been reported (Wrensch et al, 1990). We attribute this greater success to differences in technique, as well as to persistence in attempting to obtain NAF. Quite a few modifications in technique have been made since the first 11 subjects provided NAF, when the success rate was 43%. We have found that it is important to warm and massage the breast before aspiration. Allowing the subject to massage her own breast avoids compression to the point of discomfort. We aspirated each breast twice at each visit, whereas in most other reports each breast was aspirated only once. Each subject was asked to return for further visits. Early in our experience, we examined the NAF cytology in 11 subjects who underwent nipple aspiration two or three times. The NAF was collected in separate containers at each visit and cytological review was performed. The cytological classification of the specimens (Table 1) obtained from the second and third visits was the same as the first visit in 10 of 11 cases.

We felt that it was important to study subjects of a single risk category (precancerous mastopathy) when we were assessing the feasibility of the procedure. Thus, the first 33 subjects recruited all had a diagnosis of DCIS, LCIS or AH. Early into this pilot study we realized, however, that only the untreated breast in subjects with a history of DCIS could be aspirated. Thus, the pilot included subjects now classified as having either precancerous mastopathy or a personal history of breast cancer. Nine of the ten subjects in whom nipple aspiration was unsuccessful were enrolled during the pilot study. We later broadened our entry criteria to include subjects of all risk categories. Since we have standardized our aspiration technique, we have been equally successful in obtaining NAF from subjects of all risk categories.

Wrensch et al (1993) evaluated NAF in a cohort of subjects with normal breast cancer risk. In this population, they demonstrated that subjects with NAF that contained normal cytology, hyperplasia without atypia, or atypical hyperplasia have a risk of breast cancer similar to subjects who have a biopsy with similar diagnoses. They also reported that subjects with scant cellularity had the lowest risk. Our goal was to determine the usefulness of NAF cytology in subjects of all risk categories. We therefore evaluated NAF in women of each risk category, and correlated the cytological findings with breast cancer risk, based on family history or prior pathology. NAF cytology was highly correlated ($P = 0.002$, Table 4) with breast cancer risk, suggesting that cytology may be a useful biomarker for subjects of various risk categories.

Computerized image analysis is able to quantitate the DNA index and cell cycle parameters on a cell by cell basis. The prognostic utility of this modality has been demonstrated for a variety of tumours, including breast cancer (Dressler et al, 1988). Although flow cytometry is the standard method to determine cellular DNA, image analysis using Feulgen-stained cell preparations is gaining wider acceptance because of its ability to evaluate samples of relatively scant cellularity. Moreover, studies have demonstrated a good correlation between DNA indices determined by flow and image cytometry (Ellison et al, 1993).

DNA indices in epithelial cells from nipple aspirate specimens were determined and evaluated for their correlation with cytological class. Increasing DNA index, increasing percentage of cells in G₂M and increasing percentage of hyperdiploid cells were found as the cytology became more abnormal, suggesting their potential usefulness as biomarkers.

The correlation between cytology and risk (Table 4) may underestimate the true significance of the relationship because of the way in which the NAF samples were collected. Over 90% of the NAF specimens with recently diagnosed DCIS or invasive breast cancer came from a mastectomy specimen. Thus, only one aspiration attempt was possible. On the other hand, the majority of women with a normal breast cancer risk or a family history of breast cancer agreed to undergo multiple aspirations, some as many as 12. One would expect that increasing the number of samples would increase the chance of finding abnormal cells should they exist. Of the 13 subjects with normal breast cancer risk who underwent aspiration 1-6 times, no cytology specimen contained atypical cells. Of the nine subjects at normal risk who underwent aspiration 12 times, two (22%) had atypical hyperplasia (AH). The study by Wrench et al (1993) would suggest that the subjects thought to be at normal risk, with NAF cytology demonstrating AH, have a risk of breast cancer similar to if they had undergone a biopsy demonstrating atypical hyperplasia. In the two subjects with AH found in NAF, who were presumed to have normal breast cancer risk, NAF was useful in identifying that these women were at increased risk. Although a statistical correlation was not found between cytological class and the number of aspirations, it is logical that increasing the number of aspirations will increase the likelihood of finding abnormal cells. In order to increase the likelihood of finding abnormal cells, NAF may be most useful when performed more than once or at intervals, as are breast examinations or mammograms.

All NAF samples that are fixed and prepared properly are evaluable for cytology, for even a cytological classification of 1 (inadequate cells) provides important prognostic information. (Subjects with class 1 cytology seem to have the lowest breast cancer risk (Wrench et al, 1993).) As indicated in Table 3, we elected to use the entire sample from seven subjects for non-cellular studies, because of our interest in certain non-cellular markers. One of the markers, prostate-specific antigen (PSA), has proven to be a very promising marker of breast cancer risk (Sauter et al, 1996). Non-cellular markers in NAF are in theory always evaluable, so long as the protein of interest can be detected. In our hands, 1 µl of NAF is more than sufficient to detect and quantify PSA.

On the other hand, we are presently able to obtain samples of adequate cellularity for DNA ploidy and cell cycle determinations in just over 50% of subjects. Although we do not know if we can increase the percentage of subjects providing cellular samples, we feel that we can increase the quantity of breast epithelial cells obtained from subjects who do provide cellular samples. The

quantity of cells obtained is important, as a large number of cells are required for a variety of tests (e.g. proliferation score). At present, we are performing two aspirations on each breast at each visit. It is our impression that two aspirations on each breast do not remove all of the fluid from the breast. A simple method, then, to increase the number of breast epithelial cells obtained would be to perform a third aspiration on each breast at each visit. We feel that a third aspiration would be well tolerated, as the procedure is quick and causes minimal to no discomfort. Nonetheless, because not all of the specimens will contain sufficient cells for evaluation, we believe that non-cellular (e.g. PSA) as well as cellular markers of breast cancer risk should be used in combination to maximize the information obtained from the NAF sample.

We do not plan to perform nipple aspiration in a previously irradiated breast. The low yield of both NAF and cellular specimens indicates that radiation treatment significantly alters the secretion of both fluid and cells into the breast ductal lumen. We have elected not to aspirate individuals who have received chemotherapy for any malignancy because of the possible effect of this treatment on the aspiration specimen.

We can envisage two possible uses for the nipple aspirate fluid: one as an adjunct to screening and a second to evaluate intermediate markers for change in response to a chemopreventive agent. A subject coming in for her annual examination by her gynaecologist or family physician could undergo nipple aspiration either just before or just after the physical examination. The aspiration need not be done by a physician, so long as the individual is properly trained to perform the procedure. The aspiration takes only 5-10 min to perform, and can be repeated as early as 3 days later if more fluid is needed (probably even sooner, although we have not tested this). One or more of the capillary tubes containing NAF could be sent to a reference laboratory for analysis of one or more non-cellular markers, such as PSA. A second capillary tube containing NAF could be emptied into a cytological fixative, and sent to a cytologist with expertise in reading aspirates from body cavities, breast cysts, cerebrospinal fluid, etc. One microlitre is usually sufficient for both non-cellular and cellular studies, although a larger quantity is helpful. In the ideal scenario, if the technique were to become widely accepted, a buffer for the non-cellular markers and the cytological fixative for whole-cell studies, such as cytology and DNA image analysis, contained in a commercially available kit would streamline and standardize the testing of NAF. Even without this 'kit', however, the preparation of the sample for analysis is extremely simple, requiring only that one flush the NAF sample into the buffer or fixative with a 21-gauge needle. A similar approach to the collection of and preparation of NAF could be performed in subjects participating in a chemoprevention trial, in which the investigator would monitor the biomarker(s) of interest for changes from abnormal towards a more normal phenotype.

In summary, nipple aspiration is a feasible method to obtain breast epithelial cells in non-lactating women. The procedure is inexpensive, quick, non-invasive, can be repeated as needed and causes minimal to no discomfort. It is extremely successful in young women, for whom mammography is often non-revealing because of the density of the breast tissue. Despite its limitations, nipple aspiration has the potential to supplement mammography and physical examination in women who yield evaluable fluid. Given that both cytology and DNA indices are promising intermediate biomarkers, as well as the identification of promising non-cellular markers such as PSA, nipple aspiration may be a useful

tool to evaluate response to treatment with a chemopreventive drug. These markers could best be validated in a randomized, prospective trial.

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What Is Metastatic Cancer?

Local, Regional, and Distant (Metastatic) Spread of Cancer

Metastatic cancer is a cancer that has spread from its primary site (the part of the body in which it developed) to other parts of the body. If cells break away from a cancerous tumor, they can travel to other areas of the body. There, they may settle and form "colony" tumors. In their new location, the cancer cells continue growing. The spread of a tumor to a new part of the body is called metastasis.

It is important to understand the difference between metastasis and local spread, because they affect a patient's prognosis (the outlook for chances of survival) and treatment options in different ways.

Local spread means that a growing cancer extends beyond the organ in which it developed, into nearby organs and tissues. For example, the cervix (lower part of the uterus or womb) is located in front of the rectum and behind the bladder. Very large cancers of the cervix may extend into the rectum or bladder.

Metastasis involves spread of cancer cells through the bloodstream, or the lymph system. The lymph system consists of lymph vessels (similar to veins except that they contain tissue waste products and immune system cells instead of blood). These lymph vessels lead to lymph nodes, bean-shaped collections of immune system cells that are important in fighting infections. Cancer cells that break off from tumors and enter the lymph vessels may be carried to lymph nodes where they may continue to grow and form metastases. Doctors sometimes call metastasis to lymph nodes near the place a cancer developed regional spread. This is to distinguish it from distant spread or distant metastasis. Distant spread generally occurs when cancer cells break off from tumors and enter the bloodstream, travel to other organs, and continue to grow into new tumors.

When cancer spreads, it is still named after the part of the body where it started. For example, if prostate cancer spreads to the bones, it is still called prostate cancer, and if breast cancer spreads to the lungs it is still breast cancer.

Recurrence and Metastasis

Recurrence is the medical term used when cancer comes back in a

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EXHIBIT 2

patient who appeared to be in remission (free of cancer) after treatment. Cancer can recur locally, in the same organ it developed in or, if that organ was removed by surgery, in a nearby remaining organ or tissue. For example, local recurrence of breast cancer means cancer started to grow again in the same breast after lumpectomy and radiation therapy or in the chest wall, skin or muscle after a mastectomy. A regional recurrence usually means cancer has come back in nearby lymph nodes or in the area that lymph nodes had been removed from. After apparently complete removal of a mouth or throat cancer, growth of cancer in lymph nodes of the neck would be considered regional recurrence. A distant recurrence involves any other part of the body not included in local or regional recurrence. After surgery for apparently localized prostate cancer, the cancer might recur distantly in bones, the liver, or the lungs.

But, recurrence is not the only time a cancer will metastasize. Some cancers are discovered to have spread to nearby lymph nodes or distant organs shortly after the cancer is first diagnosed, and before treatment is started. In some cases, a metastasis may be discovered before the primary (original) tumor is found. For example, a patient may go to the doctor because lymph nodes in the neck are growing larger. After a biopsy reveals cancer in these lymph nodes, additional examinations or x-rays may find that the cancer actually started in the throat. Even if a cancer has spread very widely throughout the body before it is discovered, the cancer's appearance under a microscope can sometimes tell doctors where it started. But, some cancers lack these microscopic clues to their origin, and it may be impossible to determine exactly where they started. This condition is called cancer of unknown primary, and is discussed further in a separate American Cancer Society document.

So, cancer can recur without metastasizing (a local recurrence) and can metastasize without recurring locally (cancer found to have metastasized when first diagnosed).

Significance of Metastasis to the Patient

Most people who die of cancer have metastases at the time of their death. And, these metastases are directly responsible for the majority of cancer deaths. Most common cancers (prostate, breast, colon, lung, for example) develop in organs that can be completely or partially removed by surgery. Even though these operations cause effects, they could cure patients if metastasis did not occur. Most of the serious consequences of these cancers occur because of spread to other parts of the body. In some cases, the most serious effect of cancer is its spread to a particularly essential part of the body, such as areas of the brain. In other cases, spread to and growth in many organs creates so many cancer cells that the body's normal metabolism is disrupted.

However, cancer can be fatal even if it does not metastasize. Cancers that start in the brain may be impossible to remove without causing severe brain damage. In fact, brain tumors rarely spread beyond the nervous system. Growth and local spread of these cancers often interfere with parts of the brain necessary for life. A tumor may become very large without ever metastasizing. It may then press against vital tissues such as nerves, the brain, or blood vessels. Leukemia can cause death by interfering with normal blood functions without the defective cells ever leaving the tissues they started in (the bone marrow and bloodstream).

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